



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/889,802

Source: Pct/09

Date Processed by STIC: 8/1/2001

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER
VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND
TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 - 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST-25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO).

Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:

<http://www.uspto.gov/web/offices/pac/checker>

PCT09

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/889,802

DATE: 08/01/2001
TIME: 18:28:48

Input Set : A:\es.txt
Output Set: N:\CRF3\08012001\I889802.raw

pg 1-3
Does Not Comply

Corrected Diskette Needed

W--> 1 SEQUENZPROTOKOLL *delete*
3 <110> APPLICANT: Kreutzer Dr., Roland
4 Limmer Dr., Stephan
6 <120> TITLE OF INVENTION: Verfahren und Medikament zur Hemmung der Expression
7 eines vorgegebenen Gens
9 <130> FILE REFERENCE: 400968
C--> 11 <140> CURRENT APPLICATION NUMBER: US/09/889,802
C--> 12 <141> CURRENT FILING DATE: 2001-07-20
14 <150> PRIOR APPLICATION NUMBER: 199 03 713.2
15 <151> PRIOR FILING DATE: 1999-01-30
17 <150> PRIOR APPLICATION NUMBER: 199 56 568.6
18 <151> PRIOR FILING DATE: 1999-11-24
20 <160> NUMBER OF SEQ ID NOS: 8
22 <170> SOFTWARE: PatentIn Ver. 2.1
24 <210> SEQ ID NO: 1
25 <211> LENGTH: 45
26 <212> TYPE: DNA
27 <213> ORGANISM: Knstliche Sequenz
29 <220> FEATURE:
30 <223> OTHER INFORMATION: Beschreibung der knstlichen Sequenz:
31 EcoRI-Schnittstelle, T7-RNA-Polymerasepromotor
33 <400> SEQUENCE: 1
34 ggaattctaa tacgactcac tatagggcga tcagatctct agaag 45
37 <210> SEQ ID NO: 2
38 <211> LENGTH: 50
39 <212> TYPE: DNA
40 <213> ORGANISM: Knstliche Sequenz
42 <220> FEATURE:
43 <223> OTHER INFORMATION: Beschreibung der knstlichen Sequenz:
44 BamHI-Schnittstelle, SP6-RNA-Polymerasepromotor
46 <400> SEQUENCE: 2
47 gggatccatt taggtgacac tatagaatac ccatgatcgc gtagtcgata 50
50 <210> SEQ ID NO: 3
51 <211> LENGTH: 340
52 <212> TYPE: RNA
53 <213> ORGANISM: Knstliche Sequenz
55 <220> FEATURE:
56 <223> OTHER INFORMATION: Beschreibung der knstlichen Sequenz: RNA, die
57 einer Sequenz aus der "positive control DNA" des
58 HeLaScribe Nuclear Extract in vitro
59 Transkriptionskits der Firma Promega entspricht
61 <400> SEQUENCE: 3
62 ucagaucucu agaagcuuua augcgguagu uuaucaacagu uaaaauugcua acgcagucag 60
63 gcaccgugua ugaaaucuaa caaugcguc aucgucaucc ucggcaccgu caccucggau 120
64 gcuguaggca uaggcuuggu uaugccggua cugccgggcc ucuugcgga uaucguccau 180
65 uccgacgca ucgcaguca cuauggcgug cugcuagcgc uauaugcguu gaugcauuu 240
66 cuaugcgcac ccguucucgg agcacugucc gaccgcuuug gccgccgcc aguccugcuc 300

*all U.S. applications
must be in English.*

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/889,802

DATE: 08/01/2001

TIME: 18:28:48

Input Set : A:\es.txt

Output Set: N:\CRF3\08012001\I889802.raw

```

67 gcuucgcuac uuggagccac uaucgacuac gcgaucagg 340
70 <210> SEQ ID NO: 4
71 <211> LENGTH: 363
72 <212> TYPE: DNA
73 <213> ORGANISM: Knstliche Sequenz
75 <220> FEATURE:
76 <223> OTHER INFORMATION: Beschreibung der knstlichen Sequenz: DNA, die
77 einer Sequenz aus der "positive control DNA" des
78 HeLaScribe Nuclear Extract in vitro
79 Transkriptionskits der Firma Promega entspricht
81 <400> SEQUENCE: 4
82 tcagatctct agaagcttta atgcggtagt ttatcacagt taaattgcta acgcagtcag 60
83 gcaccgtgta tgaaatctaa caatgcgctc atcgtcatcc tcggcaccgt caccctggat 120
84 gctgtaggca taggcttggg tatgccgta ctgccggcc tcttgcgga tatcgtccat 180
85 tccgacagca tcgccagtca ctatggcgtg ctgctagcgc tatatgcgtt gatgcaattt 240
86 ctatgcgcac ccgttctcgg agcactgtcc gaccgcttg gccgcccgc agtctgtctc 300
87 gcttcgtac ttggagccac tatcgactac gcgatcatgg cgaccacacc cgtcctgtgg 360
88 atc 363
91 <210> SEQ ID NO: 5
92 <211> LENGTH: 315
93 <212> TYPE: RNA
94 <213> ORGANISM: Knstliche Sequenz
96 <220> FEATURE:
97 <223> OTHER INFORMATION: Beschreibung der knstlichen Sequenz: Sequenz aus
98 dem YFP-Gen
100 <400> SEQUENCE: 5
101 auggugagca agggcgagga gcguucacc gggguggugc ccauccuggu cgagcuggac 60
102 ggcgacguaa acggccacaa guucagcgug uccggcgagg gcgagggcga ugccaccuac 120
103 ggcaagcuga ccugaaguu caucugcacc accggcaagc ugcccugugc cuggcccacc 180
104 cucgugacca ccugaccua cggcgugcag ugcuuacagc gcuaccccga ccacaugaag 240
105 cagcacgacu ucuucaaguc cgccaugccc gaaggcuacg uccaggagcg caccaucuuc 300
106 uucaaggacg acggc 315
109 <210> SEQ ID NO: 6
110 <211> LENGTH: 52
111 <212> TYPE: DNA
112 <213> ORGANISM: Knstliche Sequenz
114 <220> FEATURE:
115 <223> OTHER INFORMATION: Beschreibung der knstlichen Sequenz:
116 EcoRI-Schnittstelle, T7-RNA-Polymerasepromotor,
117 komplementärer Bereich zum YFP-Gen
119 <400> SEQUENCE: 6
120 ggaattctaa tacgactcac tatagggcga atggtgagca agggcgagga gc 52
123 <210> SEQ ID NO: 7
124 <211> LENGTH: 53
125 <212> TYPE: DNA
126 <213> ORGANISM: Knstliche Sequenz
128 <220> FEATURE:
129 <223> OTHER INFORMATION: Beschreibung der knstlichen Sequenz:
130 BamHI-Schnittstelle, SP6-RNA-Polymerasepromotor,

```

RAW SEQUENCE LISTING

DATE: 08/01/2001

PATENT APPLICATION: US/09/889,802

TIME: 18:28:48

Input Set : A:\es.txt

Output Set: N:\CRF3\08012001\I889802.raw

131 komplement, rer Bereich zum YFP-Gen
133 <400> SEQUENCE: 7
134 gggatccatt taggtgacac tatagaatac gccgtcgtcc ttgaagaaga tgg 53
137 <210> SEQ ID NO: 8
138 <211> LENGTH: 21
139 <212> TYPE: RNA
140 <213> ORGANISM: Knstliche Sequenz
142 <220> FEATURE:
143 <223> OTHER INFORMATION: Beschreibung der knstlichen Sequenz: RNA, die
144 einer Sequenz aus dem YFP-Gen entspricht
146 <400> SEQUENCE: 8
147 ucgagcugga cggcgacgua a 21

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/889,802

DATE: 08/01/2001

TIME: 18:28:49

Input Set : A:\es.txt

Output Set: N:\CRF3\08012001\I889802.raw

L:1 M:259 W: Allowed number of lines exceeded, (1) GENERAL INFORMATION:
L:11 M:270 C: Current Application Number differs, Replaced Application Number
L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date

STATISTICS SUMMARY

PATENT APPLICATION: US/09/889,802

DATE: 08/01/2001

TIME: 18:28:49

Input Set : A:\es.txt

Output Set: N:\CRF3\08012001\I889802.raw

Application Serial Number: US/09/889,802

Alpha or Numeric: Numeric

Application Class:

Application File Date: 07-20-2001

Art Unit: PCT09

Software Application: PatentIn

Total Number of Sequences: 8

Total Nucleotides: 1239

Total Amino Acids: 0

Number of Errors: 0

Number of Warnings: 1

Number of Corrections: 2

MESSAGE SUMMARY

259 W: 1 (Allowed number of lines exceeded)

270 C: 1 (Current Application Number differs)

271 C: 1 (Current Filing Date differs)